

**Amendments to the Specification:**

Please replace the paragraph which begins at page 29, line 35, and ends at page 30, line 14, with the following amended paragraph:

Figure 3 (a) to (k) - This shows the nucleotide sequences of CryIA (b&c), the ricin toxin B chain gene fragments and the fusion genes cloned in pFASTBAC1 under the control of the polyhedrin promoter. The sequences in figures 3(a)-(k) are labeled Seq ID Nos 1-11, respectively. All sequence read 5'-3' direction. The codon ATG starting at nucleotide 97 is the translation initiation site for the genes CryIA(b&c) and all the fusion genes. For the ricin toxin B chain fragments the codon ATG starting at nucleotide position 125 serve as the translation initiation site. All the genes are terminated by the SV40 polyadenylation sequence. The stop codons TAG and TAA are employed and their positions (if sequences are read in the 5'-3'direction) vary with the size of each gene. If the sequences are read from the 3'-5' direction the two codons TAG and TAA are located at nucleotide

positions 17 and 7 respectively.

Please replace the paragraph at page 31, lines 9-12, with the following amended paragraph:

LF1=5' CAACAACAAAGGAATTCATGCTGATG 3' (Seq ID No 12)

LB1=5' GGACACACACACTGCAAGCTTGTAATC 3' (Seq ID No 13)

LB2=5' CGGATCCGAAAGCTTCACATCTAACAC 3' (Seq ID No 14)

LB3=5' GCTTGCAAGCTTAGACCATATAGCCC 3' (Seq ID No 15)

Please replace the paragraph begins at page 43, line 34, ends at page 44, line 9, with the following amended paragraph:

Total RNA was extracted from 100 mg leaf tissue of transformed and wild type rice plants using the Rneasy Plant Mini kit (Qiagen) according to the supplier's recommendations. RT-PCRs were carried out using the Access-PCR kit (Promega) according to the manufacturer's instructions. We used 100 ng total RNA and 50 pmol of each primer. Primers CRF1 and CRR1 amplify both *cry1Ab* and *cry1Ac*, while primers RTF1 and RTR1 amplify the RTB gene fragment. The primer sequences were as follows: CRF1 (5'-

CGCATTGAAAC CGGTTACACTC CCA-3' (Seq ID No 16)), CRR1 (5'-CTTGGGCAGAACCACGGAAGCTACC-3' (Seq ID No 17)), RTF1 (5'-GATGTTTGTATGGATCCTCAGCCCA-3' (Seq ID No 18)) and RTR1 (5'-GCCGAACAATGGTTGTAACAAAAGG-3' (Seq ID No 19)).